

Fig. 1

1 TCAATCAACGAGGCGTCCGTGCGACACAGGAGGAAATCCAATGAGCGGAAAACTGGCTTA 60
M S G K L A Y

61 CGTTACAGGCGGGATGGGCGGTATCGGCACCTCAATTTGCCAGCGCCTGGCCAAAGATGG 120
V T G G M G G I G T S I C Q R L A K D G

121 CTTTCGCGTGGTGGCAGGCTGCGGCCCCAGCCGCAATTACCAGCAATGGCTGGATGAACA 180
F R V V A G C G P S R N Y Q Q W L D E Q

181 GGC GGCGCAGGGCTATACGTTCTACGCGTCAGTGGGCAACGTGTCCGATTGGGAGTCCAC 240
A A Q G Y T F Y A S V G N V S D W E S T

241 GGTAGAAGCATTGAGCGCGTCAAGCGGGACATGGGCCCGGTGATGTGCTGGTCAACAA 300
V E A F E R V K R D M G P V D V L V N N

301 CGCGGGCATCACCCGCGACGGCCTGTTCCGCAAGATGAGCGCCGACGACTGGCGCGCGGT 360
A G I T R D G L F R K M S A D D W R A V

361 CATCGACACCAACCTGAACAGCCTCTTCAACGTGACCAAGCAGGTGATCGACGACATGGT 420
I D T N L N S L F N V T K Q V I D D M V

421 CGAGCGCCAGTGGGGCCGCATCGTCAACATCAGCTCGGTGAACGGGCAGAAGGGGCAGTT 480
E R Q W G R I V N I S S V N G Q K G Q F

481 CGGCCAGACGAACTATTCACGGCGAAGGCGGGCATCCATGGCTTCACCATGGCGCTGGC 540
G Q T N Y S T A K A G I H G F T M A L A

541 GCAGGAAGTGGCCAGCAAGGGCATCACGGTCAACACGGTGTGCGCGGGCTACATCGGCAC 600
Q E V A S K G I T V N T V S P G Y I G T

601 GGACATGGTTCGCGCCATCCGTCCGGACGTGCTGGAAAAGATCGTCGCCACCATTCCGGT 660
D M V R A I R P D V L E K I V A T I P V

661 GCGCCGCTGGGCACGCCGGAGGAAATCGCGTCCATCACGTGCTGGCTGGCCTCGGATGA 720
R R L G T P E E I A S I T S W L A S D E

721 GTCTGGGTTTTGACGGGCGCGGACTTCTCGCTCAACGGCGGCCTGCATATGGGCTGAAG 780
S G F S T G A D F S L N G G L H M G *

781 CATCGCGGGCCGCCACGAGCGGCCCCGCGCGCGGGCGGCCTCGGGGAGAGGGCCGTCC 840

841 GGCATTACACTTACCCTTATCCGAAGTCTTAGAGATCGCCCGATCCGGGGACAACCATGA 900

Fig. 2

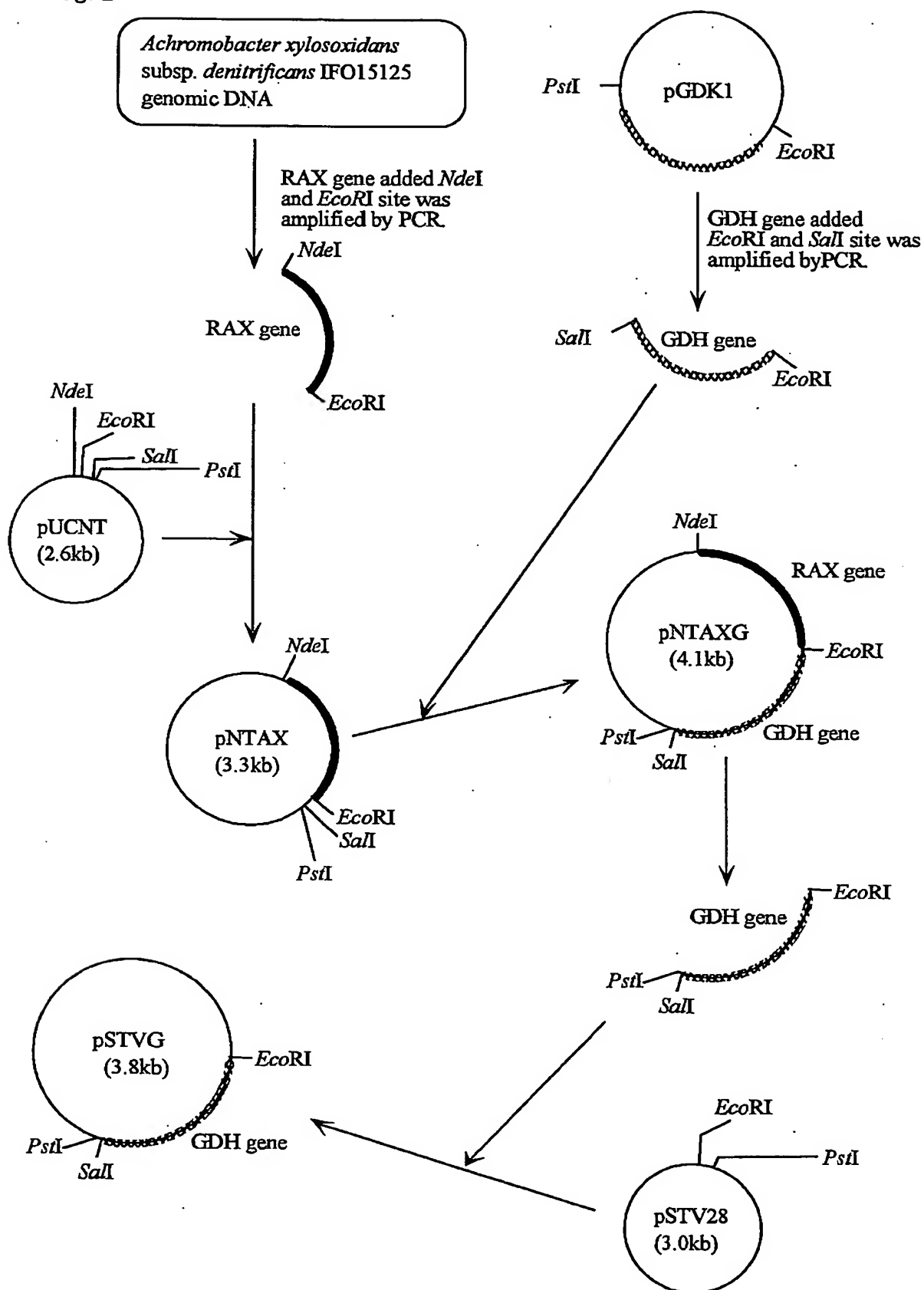


Fig. 3

